

SEQUENCE LISTING

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<120> COMPOSITIONS AND METHODS FOR MODIFICATION AND PREVENTION OF SARS
CORONAVIRUS INFECTIVITY

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Ala Tyr 1070	Phe Pro Arg Glu Gly 1075	Val Phe Val Phe Asn 1080
Trp Phe 1085	Ile Thr Gln Arg Asn 1090	Phe Phe Ser Pro Gln 1095
Thr Asp 1100	Asn Thr Phe Val Ser 1105	Gly Asn Cys Asp Val 1110
Ile Ile 1115	Asn Asn Thr Val Tyr 1120	Asp Pro Leu Gln Pro 1125
Ser Phe 1130	Lys Glu Glu Leu Asp 1135	Lys Tyr Phe Lys Asn 1140
Pro Asp 1145	Val Asp Leu Gly Asp 1150	Ile Ser Gly Ile Asn 1155
Val Asn 1160	Ile Gln Lys Glu Ile 1165	Asp Arg Leu Asn Glu 1170
Asn Leu 1175	Asn Glu Ser Leu Ile 1180	Asp Leu Gln Glu Leu 1185
Glu Gln 1190	Tyr Ile Lys Trp Pro 1195	Trp Tyr Val Trp Leu 1200
Ala Gly 1205	Leu Ile Ala Ile Val 1210	Met Val Thr Ile Leu 1215
Met Thr 1220	Ser Cys Cys Ser Cys 1225	Leu Lys Gly Ala Cys 1230
Ser Cys 1235	Cys Lys Phe Asp Glu 1240	Asp Asp Ser Glu Pro 1245
Gly Val 1250	Lys Leu His Tyr Thr 1255	

<210> 3
 <211> 390
 <212> DNA
 <213> SARS coronavirus Urbani

<220>
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 <222> (1)..(390)

<400> 3
atg caa atg gca tat agg ttc aat ggc att gga gtt acc caa aat gtt 48
Met Gln Met Ala Tyr Arg Phe Asn Gly Ile Gly Val Thr Gln Asn Val
1 5 10 15

ctc tat gag aac caa aaa caa atc gcc aac caa ttt aac aag gcg att 96
Leu Tyr Glu Asn Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile
20 25 30

agt caa att caa gaa tca ctt aca aca aca tca act gca ttg ggc aag 144
Ser Gln Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys
35 40 45

ctg caa gac gtt gtt aac cag aat gct caa gca tta aac aca ctt gtt 192
Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val
50 55 60

aaa caa ctt agc tct aat ttt ggt gca att tca agt gtg cta aat gat 240
Lys Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn Asp
65 70 75 80

atc ctt tcg cga ctt gat aaa gtc gag gcg gag gta caa att gac agg 288
Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val Gln Ile Asp Arg
85 90 95

tta att aca ggc aga ctt caa agc ctt caa acc tat gta aca caa caa 336
Leu Ile Thr Gly Arg Leu Gln Ser Leu Gln Thr Tyr Val Thr Gln Gln
100 105 110

cta atc agg gct gct gaa atc agg gct tct gct aat ctt gct gct act 384
Leu Ile Arg Ala Ala Glu Ile Arg Ala Ser Ala Asn Leu Ala Ala Thr
115 120 125

aaa atg 390
Lys Met
130

<210> 4
<211> 130
<212> PRT
<213> SARS coronavirus Urbani

<400> 4
Met Gln Met Ala Tyr Arg Phe Asn Gly Ile Gly val Thr Gln Asn val
1 5 10 15

Leu Tyr Glu Asn Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile
20 25 30

ser Gln Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys
35 40 45

Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val
50 55 60

Lys Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser val Leu Asn Asp
65 70 75 80

Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu val Gln Ile Asp Arg
85 90 95

Leu Ile Thr Gly Arg Leu Gln Ser Leu Gln Thr Tyr Val Thr Gln Gln
100 105 110

Leu Ile Arg Ala Ala Glu Ile Arg Ala Ser Ala Asn Leu Ala Ala Thr
115 120 125

Lys Met
130

<210> 5
<211> 276
<212> DNA
<213> SARS coronavirus Urbani

<220>
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<222> (1)..(276)

<400> 5
atg caa atg gca tat agg ttc aat ggc att gga gtt acc caa aat gtt 48
Met Gln Met Ala Tyr Arg Phe Asn Gly Ile Gly Val Thr Gln Asn Val
1 5 10 15
ctc tat gag aac caa aaa caa atc gcc aac caa ttt aac aag gcg att 96
Leu Tyr Glu Asn Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile
20 25 30
agt caa att caa gaa tca ctt aca aca aca tca act gca ttg ggc aag 144
Ser Gln Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys
35 40 45
ctg caa gac gtt gtt aac cag aat gct caa gca tta aac aca ctt gtt 192
Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val
50 55 60
aaa caa ctt agc tct aat ttt ggt gca att tca agt gtg cta aat gat 240
Lys Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn Asp
65 70 75 80
atc ctt tcg cga ctt gat aaa gtc gag gcg gag gta 276
Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val
85 90

<210> 6
<211> 92
<212> PRT
<213> SARS coronavirus Urbani

<400> 6
Met Gln Met Ala Tyr Arg Phe Asn Gly Ile Gly Val Thr Gln Asn Val
1 5 10 15
Leu Tyr Glu Asn Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile
20 25 30
Ser Gln Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys
35 40 45
Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val
50 55 60

Lys Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn Asp
65 70 75 80

Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val
85 90

<210> 7
<211> 174
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(174)

<400> 7
att caa gaa tca ctt aca aca aca tca act gca ttg ggc aag ctg caa 48
Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15
gac gtt gtt aac cag aat gct caa gca tta aac aca ctt gtt aaa caa 96
Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln
20 25 30
ctt agc tct aat ttt ggt gca att tca agt gtg cta aat gat atc ctt 144
Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn Asp Ile Leu
35 40 45
tcg cga ctt gat aaa gtc gag gcg gag gta 174
Ser Arg Leu Asp Lys Val Glu Ala Glu Val
50 55

<210> 8
<211> 58
<212> PRT
<213> SARS coronavirus Urbani

<400> 8
Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15
Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln
20 25 30
Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn Asp Ile Leu
35 40 45
Ser Arg Leu Asp Lys Val Glu Ala Glu Val
50 55

<210> 9
<211> 141
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS

<222> (1)..(141)

<400> 9

ttg ggc aag ctg caa gac gtt gtt aac cag aat gct caa gca tta aac	48
Leu Gly Lys Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn	
1 5 10 15	

aca ctt gtt aaa caa ctt agc tct aat ttt ggt gca att tca agt gtg	96
Thr Leu Val Lys Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val	
20 25 30	

cta aat gat atc ctt tcg cga ctt gat aaa gtc gag gcg gag gta	141
Leu Asn Asp Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val	
35 40 45	

<210> 10

<211> 47

<212> PRT

<213> SARS coronavirus Urbani

<400> 10

Leu Gly Lys Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn	
1 5 10 15	

Thr Leu Val Lys Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val	
20 25 30	

Leu Asn Asp Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val	
35 40 45	

<210> 11

<211> 114

<212> DNA

<213> SARS coronavirus Urbani

<220>

<221> CDS

<222> (1)..(114)

<400> 11

caa att gac agg tta att aca ggc aga ctt caa agc ctt caa acc tat	48
Gln Ile Asp Arg Leu Ile Thr Gly Arg Leu Gln Ser Leu Gln Thr Tyr	
1 5 10 15	

gta aca caa caa cta atc agg gct gct gaa atc agg gct tct gct aat	96
Val Thr Gln Gln Leu Ile Arg Ala Ala Glu Ile Arg Ala Ser Ala Asn	
20 25 30	

ctt gct gct act aaa atg	114
Leu Ala Ala Thr Lys Met	
35	

<210> 12

<211> 38

<212> PRT

<213> SARS coronavirus Urbani

<400> 12

Gln Ile Asp Arg Leu Ile Thr Gly Arg Leu Gln Ser Leu Gln Thr Tyr	
1 5 10 15	

Val Thr Gln Gln Leu Ile Arg Ala Ala Glu Ile Arg Ala Ser Ala Asn
20 25 30

Leu Ala Ala Thr Lys Met
35

<210> 13
<211> 105
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(105)

<400> 13
atg caa atg gca tat agg ttc aat ggc att gga gtt acc caa aat gtt 48
Met Gln Met Ala Tyr Arg Phe Asn Gly Ile Gly Val Thr Gln Asn Val
1 5 10 15
ctc tat gag aac caa aaa caa atc gcc aac caa ttt aac aag gcg att 96
Leu Tyr Glu Asn Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile
20 25 30
agt caa att 105
Ser Gln Ile
35

<210> 14
<211> 35
<212> PRT
<213> SARS coronavirus Urbani

<400> 14
Met Gln Met Ala Tyr Arg Phe Asn Gly Ile Gly Val Thr Gln Asn Val
1 5 10 15
Leu Tyr Glu Asn Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile
20 25 30
Ser Gln Ile
35

<210> 15
<211> 105
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(105)

<400> 15
ttc aat ggc att gga gtt acc caa aat gtt ctc tat gag aac caa aaa 48
Phe Asn Gly Ile Gly Val Thr Gln Asn Val Leu Tyr Glu Asn Gln Lys
1 5 10 15
caa atc gcc aac caa ttt aac aag gcg att agt caa att caa gaa tca 96
Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile Ser Gln Ile Gln Glu Ser

	20	25	30	105
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ctt aca aca
 Leu Thr Thr
 35

<210> 16
 <211> 35
 <212> PRT
 <213> SARS coronavirus Urbani
 <400> 16

Phe Asn Gly Ile Gly Val Thr Gln Asn Val Leu Tyr Glu Asn Gln Lys
 1 5 10 15

Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile Ser Gln Ile Gln Glu Ser
 20 25 30

Leu Thr Thr
 35

<210> 17
 <211> 105
 <212> DNA
 <213> SARS coronavirus Urbani
 <220>
 <221> CDS
 <222> (1)..(105)
 <400> 17

caa aat gtt ctc tat gag aac caa aaa caa atc gcc aac caa ttt aac	48
Gln Asn Val Leu Tyr Glu Asn Gln Lys Gln Ile Ala Asn Gln Phe Asn	
1 5 10 15	

aag gcg att agt caa att caa gaa tca ctt aca aca aca tca act gca	96
Lys Ala Ile Ser Gln Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala	
20 25 30	

ttg ggc aag
 Leu Gly Lys
 35

<210> 18
 <211> 35
 <212> PRT
 <213> SARS coronavirus Urbani
 <400> 18

Gln Asn Val Leu Tyr Glu Asn Gln Lys Gln Ile Ala Asn Gln Phe Asn
 1 5 10 15

Lys Ala Ile Ser Gln Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala
 20 25 30

Leu Gly Lys
 35

<210> 19
 <211> 105
 <212> DNA
 <213> SARS coronavirus Urbani

<220>
 <221> CDS
 <222> (1)..(105)

<400> 19
 caa aaa caa atc gcc aac caa ttt aac aag gcg att agt caa att caa 48
 Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile Ser Gln Ile Gln
 1 5 10 15
 gaa tca ctt aca aca aca tca act gca ttg ggc aag ctg caa gac gtt 96
 Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln Asp Val
 20 25 30
 gtt aac cag 105
 Val Asn Gln
 35

<210> 20
 <211> 35
 <212> PRT
 <213> SARS coronavirus Urbani

<400> 20
 Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile Ser Gln Ile Gln
 1 5 10 15
 Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln Asp Val
 20 25 30
 Val Asn Gln
 35

<210> 21
 <211> 105
 <212> DNA
 <213> SARS coronavirus Urbani

<220>
 <221> CDS
 <222> (1)..(105)

<400> 21
 ttt aac aag gcg att agt caa att caa gaa tca ctt aca aca aca tca 48
 Phe Asn Lys Ala Ile Ser Gln Ile Gln Glu Ser Leu Thr Thr Thr Ser
 1 5 10 15
 act gca ttg ggc aag ctg caa gac gtt gtt aac cag aat gct caa gca 96
 Thr Ala Leu Gly Lys Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala
 20 25 30
 tta aac aca 105
 Leu Asn Thr
 35

<210> 22
 <211> 35

<212> PRT
<213> SARS coronavirus Urbani

<400> 22

Phe Asn Lys Ala Ile Ser Gln Ile Gln Glu Ser Leu Thr Thr Thr Ser
1 5 10 15

Thr Ala Leu Gly Lys Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala
20 25 30

Leu Asn Thr
35

<210> 23
<211> 105
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(105)

<400> 23
att caa gaa tca ctt aca aca aca tca act gca ttg ggc aag ctg caa 48
Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15

gac gtt gtt aac cag aat gct caa gca tta aac aca ctt gtt aaa caa 96
Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln
20 25 30

ctt agc tct 105
Leu Ser Ser
35

<210> 24
<211> 35
<212> PRT
<213> SARS coronavirus Urbani

<400> 24

Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15

Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln
20 25 30

Leu Ser Ser
35

<210> 25
<211> 105
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(105)

<400> 25
aca tca act gca ttg ggc aag ctg caa gac gtt gtt aac cag aat gct 48
Thr Ser Thr Ala Leu Gly Lys Leu Gln Asp Val Val Asn Gln Asn Ala
1 5 10 15
caa gca tta aac aca ctt gtt aaa caa ctt agc tct aat ttt ggt gca 96
Gln Ala Leu Asn Thr Leu Val Lys Gln Leu Ser Ser Asn Phe Gly Ala
20 25 30
att tca agt 105
Ile Ser Ser
35

<210> 26
<211> 35
<212> PRT
<213> SARS coronavirus Urbani

<400> 26
Thr Ser Thr Ala Leu Gly Lys Leu Gln Asp Val Val Asn Gln Asn Ala
1 5 10 15
Gln Ala Leu Asn Thr Leu Val Lys Gln Leu Ser Ser Asn Phe Gly Ala
20 25 30
Ile Ser Ser
35

<210> 27
<211> 105
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(105)

<400> 27
caa gac gtt gtt aac cag aat gct caa gca tta aac aca ctt gtt aaa 48
Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val Lys
1 5 10 15
caa ctt agc tct aat ttt ggt gca att tca agt gtg cta aat gat atc 96
Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn Asp Ile
20 25 30
ctt tcg cga 105
Leu Ser Arg
35

<210> 28
<211> 35
<212> PRT
<213> SARS coronavirus Urbani

<400> 28
Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val Lys
1 5 10 15

Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn Asp Ile
20 25 30

Leu Ser Arg
35

<210> 29
<211> 105
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(105)

<400> 29
gct caa gca tta aac aca ctt gtt aaa caa ctt agc tct aat ttt ggt 48
Ala Gln Ala Leu Asn Thr Leu Val Lys Gln Leu Ser Ser Asn Phe Gly
1 5 10 15
gca att tca agt gtg cta aat gat atc ctt tcg cga ctt gat aaa gtc 96
Ala Ile Ser Ser Val Leu Asn Asp Ile Leu Ser Arg Leu Asp Lys Val
20 25 30
gag gcg gag 105
Glu Ala Glu
35

<210> 30
<211> 35
<212> PRT
<213> SARS coronavirus Urbani

<400> 30
Ala Gln Ala Leu Asn Thr Leu Val Lys Gln Leu Ser Ser Asn Phe Gly
1 5 10 15
Ala Ile Ser Ser Val Leu Asn Asp Ile Leu Ser Arg Leu Asp Lys Val
20 25 30
Glu Ala Glu
35

<210> 31
<211> 105
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(105)

<400> 31
gtt aaa caa ctt agc tct aat ttt ggt gca att tca agt gtg cta aat 48
Val Lys Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn
1 5 10 15
gat atc ctt tcg cga ctt gat aaa gtc gag gcg gag gta caa att gac 96
Asp Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val Gln Ile Asp
20 25 30

agg tta att
Arg Leu Ile
35

105

<210> 32
<211> 35
<212> PRT
<213> SARS coronavirus Urbani

<400> 32

Val Lys Gln Leu Ser Ser Asn Phe Gly Ala Ile Ser Ser Val Leu Asn
1 5 10 15

Asp Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val Gln Ile Asp
20 25 30

Arg Leu Ile
35

<210> 33
<211> 105
<212> DNA
<213> SARS coronavirus Urbani

<220>
<221> CDS
<222> (1)..(105)

<400> 33

ttt ggt gca att tca agt gtg cta aat gat atc ctt tcg cga ctt gat
Phe Gly Ala Ile Ser Ser Val Leu Asn Asp Ile Leu Ser Arg Leu Asp
1 5 10 15 48

aaa gtc gag gcg gag gta caa att gac agg tta att aca ggc aga ctt
Lys Val Glu Ala Glu Val Gln Ile Asp Arg Leu Ile Thr Gly Arg Leu
20 25 30 96

caa agc ctt
Gln Ser Leu
35 105

<210> 34
<211> 35
<212> PRT
<213> SARS coronavirus Urbani

<400> 34

Phe Gly Ala Ile Ser Ser Val Leu Asn Asp Ile Leu Ser Arg Leu Asp
1 5 10 15

Lys Val Glu Ala Glu Val Gln Ile Asp Arg Leu Ile Thr Gly Arg Leu
20 25 30

Gln Ser Leu
35

<210> 35

<211> 105
 <212> DNA
 <213> SARS coronavirus Urbani

<220>
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 <222> (1)..(105)

<400> 35
 cta aat gat atc ctt tcg cga ctt gat aaa gtc gag gcg gag gta caa 48
 Leu Asn Asp Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val Gln
 1 5 10 15
 att gac agg tta att aca ggc aga ctt caa agc ctt caa acc tat gta 96
 Ile Asp Arg Leu Ile Thr Gly Arg Leu Gln Ser Leu Gln Thr Tyr Val
 20 25 30
 aca caa caa 105
 Thr Gln Gln
 35

<210> 36
 <211> 35
 <212> PRT
 <213> SARS coronavirus Urbani

<400> 36
 Leu Asn Asp Ile Leu Ser Arg Leu Asp Lys Val Glu Ala Glu Val Gln
 1 5 10 15
 Ile Asp Arg Leu Ile Thr Gly Arg Leu Gln Ser Leu Gln Thr Tyr Val
 20 25 30
 Thr Gln Gln
 35

<210> 37
 <211> 105
 <212> DNA
 <213> SARS coronavirus Urbani

<220>
 <221> CDS
 <222> (1)..(105)

<400> 37
 ctt gat aaa gtc gag gcg gag gta caa att gac agg tta att aca ggc 48
 Leu Asp Lys Val Glu Ala Glu Val Gln Ile Asp Arg Leu Ile Thr Gly
 1 5 10 15
 aga ctt caa agc ctt caa acc tat gta aca caa caa cta atc agg gct 96
 Arg Leu Gln Ser Leu Gln Thr Tyr Val Thr Gln Gln Leu Ile Arg Ala
 20 25 30
 gct gaa atc 105
 Ala Glu Ile
 35

<210> 38
 <211> 35
 <212> PRT

<213> SARS coronavirus Urbani

<400> 38

Leu Asp Lys Val Glu Ala Glu Val Gln Ile Asp Arg Leu Ile Thr Gly
1 5 10 15

Arg Leu Gln Ser Leu Gln Thr Tyr Val Thr Gln Gln Leu Ile Arg Ala
20 25 30

Ala Glu Ile
35

<210> 39

<211> 123

<212> DNA

<213> SARS coronavirus Urbani

<220>

<221> CDS

<222> (1)..(123)

<400> 39

gat gtt gat ctt ggc gac att tca ggc att aac gct tct gtc gtc aac 48
Asp Val Asp Leu Gly Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn
1 5 10 15

att caa aaa gaa att gac cgc ctc aat gag gtc gct aaa aat tta aat 96
Ile Gln Lys Glu Ile Asp Arg Leu Asn Glu Val Ala Lys Asn Leu Asn
20 25 30

gaa tca ctc att gac ctt caa gaa ttg 123
Glu Ser Leu Ile Asp Leu Gln Glu Leu
35 40

<210> 40

<211> 41

<212> PRT

<213> SARS coronavirus Urbani

<400> 40

Asp Val Asp Leu Gly Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn
1 5 10 15

Ile Gln Lys Glu Ile Asp Arg Leu Asn Glu Val Ala Lys Asn Leu Asn
20 25 30

Glu Ser Leu Ile Asp Leu Gln Glu Leu
35 40

<210> 41

<211> 63

<212> DNA

<213> SARS coronavirus Urbani

<220>

<221> CDS

<222> (1)..(63)

<400> 41
 att gac cgc ctc aat gag gtc gct aaa aat tta aat gaa tca ctc att 48
 Ile Asp Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile
 1 5 10 15

gac ctt caa gaa ttg 63
 Asp Leu Gln Glu Leu
 20

<210> 42
 <211> 21
 <212> PRT
 <213> SARS coronavirus Urbani

<400> 42
 Ile Asp Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile
 1 5 10 15

Asp Leu Gln Glu Leu
 20

<210> 43
 <211> 84
 <212> DNA
 <213> SARS coronavirus Urbani

<220>
 <221> CDS
 <222> (1)..(84)

<400> 43
 gtc gtc aac att caa aaa gaa att gac cgc ctc aat gag gtc gct aaa 48
 Val Val Asn Ile Gln Lys Glu Ile Asp Arg Leu Asn Glu Val Ala Lys
 1 5 10 15

aat tta aat gaa tca ctc att gac ctt caa gaa ttg 84
 Asn Leu Asn Glu Ser Leu Ile Asp Leu Gln Glu Leu
 20 25

<210> 44
 <211> 28
 <212> PRT
 <213> SARS coronavirus Urbani

<400> 44
 val val Asn Ile Gln Lys Glu Ile Asp Arg Leu Asn Glu Val Ala Lys
 1 5 10 15

Asn Leu Asn Glu Ser Leu Ile Asp Leu Gln Glu Leu
 20 25

<210> 45
 <211> 105
 <212> DNA
 <213> SARS coronavirus Urbani

<220>
 <221> CDS
 <222> (1)..(105)

<400> 45
 att tca ggc att aac gct tct gtc gtc aac att caa aaa gaa att gac 48
 Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile Asp
 1 5 10 15
 cgc ctc aat gag gtc gct aaa aat tta aat gaa tca ctc att gac ctt 96
 Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp Leu
 20 25 30
 caa gaa ttg 105
 Gln Glu Leu
 35

<210> 46
 <211> 35
 <212> PRT
 <213> SARS coronavirus Urbani

<400> 46
 Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile Asp
 1 5 10 15
 Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp Leu
 20 25 30
 Gln Glu Leu
 35

<210> 47
 <211> 49
 <212> PRT
 <213> SARS coronavirus Urbani

<400> 47
 Gln Lys Gln Ile Ala Asn Gln Phe Asn Lys Ala Ile Ser Gln Ile Gln
 1 5 10 15
 Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln Asp Val
 20 25 30
 Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln Leu Ser
 35 40 45
 Ser

<210> 48
 <211> 36
 <212> PRT
 <213> SARS coronavirus Urbani

<400> 48
 Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile
 1 5 10 15
 Asp Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp

	20	25	30
Leu Gln Glu Leu			
35			
<210> 49			
<211> 1255			
<212> PRT			
<213> SARS coronavirus Urbani			
<400> 49			
Met Phe Ile Phe Leu Leu Phe Leu Thr Leu Thr Ser Gly Ser Asp Leu			
1 5 10 15			
Asp Arg Cys Thr Thr Phe Asp Asp Val Gln Ala Pro Asn Tyr Thr Gln			
20 25 30			
His Thr Ser Ser Met Arg Gly Val Tyr Tyr Pro Asp Glu Ile Phe Arg			
35 40 45			
Ser Asp Thr Leu Tyr Leu Thr Gln Asp Leu Phe Leu Pro Phe Tyr Ser			
50 55 60			
Asn Val Thr Gly Phe His Thr Ile Asn His Thr Phe Gly Asn Pro Val			
65 70 75 80			
Ile Pro Phe Lys Asp Gly Ile Tyr Phe Ala Ala Thr Glu Lys Ser Asn			
85 90 95			
Val Val Arg Gly Trp Val Phe Gly Ser Thr Met Asn Asn Lys Ser Gln			
100 105 110			
Ser Val Ile Ile Ile Asn Asn Ser Thr Asn Val Val Ile Arg Ala Cys			
115 120 125			
Asn Phe Glu Leu Cys Asp Asn Pro Phe Phe Ala Val Ser Lys Pro Met			
130 135 140			
Gly Thr Gln Thr His Thr Met Ile Phe Asp Asn Ala Phe Asn Cys Thr			
145 150 155 160			
Phe Glu Tyr Ile Ser Asp Ala Phe Ser Leu Asp Val Ser Glu Lys Ser			
165 170 175			
Gly Asn Phe Lys His Leu Arg Glu Phe Val Phe Lys Asn Lys Asp Gly			
180 185 190			
Phe Leu Tyr Val Tyr Lys Gly Tyr Gln Pro Ile Asp Val Val Arg Asp			
195 200 205			
Leu Pro Ser Gly Phe Asn Thr Leu Lys Pro Ile Phe Lys Leu Pro Leu			
210 215 220			

Gly Ile Asn Ile Thr Asn Phe Arg Ala Ile Leu Thr Ala Phe Ser Pro
 225 230 235 240
 Ala Gln Asp Ile Trp Gly Thr Ser Ala Ala Ala Tyr Phe Val Gly Tyr
 245 250 255
 Leu Lys Pro Thr Thr Phe Met Leu Lys Tyr Asp Glu Asn Gly Thr Ile
 260 265 270
 Thr Asp Ala Val Asp Cys Ser Gln Asn Pro Leu Ala Glu Leu Lys Cys
 275 280 285
 Ser Val Lys Ser Phe Glu Ile Asp Lys Gly Ile Tyr Gln Thr Ser Asn
 290 295 300
 Phe Arg Val Val Pro Ser Gly Asp Val Val Arg Phe Pro Asn Ile Thr
 305 310 315 320
 Asn Leu Cys Pro Phe Gly Glu Val Phe Asn Ala Thr Lys Phe Pro Ser
 325 330 335
 Val Tyr Ala Trp Glu Arg Lys Lys Ile Ser Asn Cys Val Ala Asp Tyr
 340 345 350
 Ser Val Leu Tyr Asn Ser Thr Phe Phe Ser Thr Phe Lys Cys Tyr Gly
 355 360 365
 Val Ser Ala Thr Lys Leu Asn Asp Leu Cys Phe Ser Asn Val Tyr Ala
 370 375 380
 Asp Ser Phe Val Val Lys Gly Asp Asp Val Arg Gln Ile Ala Pro Gly
 385 390 395 400
 Gln Thr Gly Val Ile Ala Asp Tyr Asn Tyr Lys Leu Pro Asp Asp Phe
 405 410 415
 Met Gly Cys Val Leu Ala Trp Asn Thr Arg Asn Ile Asp Ala Thr Ser
 420 425 430
 Thr Gly Asn Tyr Asn Tyr Lys Tyr Arg Tyr Leu Arg His Gly Lys Leu
 435 440 445
 Arg Pro Phe Glu Arg Asp Ile Ser Asn Val Pro Phe Ser Pro Asp Gly
 450 455 460
 Lys Pro Cys Thr Pro Pro Ala Leu Asn Cys Tyr Trp Pro Leu Asn Asp
 465 470 475 480
 Tyr Gly Phe Tyr Thr Thr Thr Gly Ile Gly Tyr Gln Pro Tyr Arg Val
 485 490 495

Val Val Leu Ser Phe Glu Leu Leu Asn Ala Pro Ala Thr Val Cys Gly
 500 505 510
 Pro Lys Leu Ser Thr Asp Leu Ile Lys Asn Gln Cys Val Asn Phe Asn
 515 520 525
 Phe Asn Gly Leu Thr Gly Thr Gly Val Leu Thr Pro Ser Ser Lys Arg
 530 535 540
 Phe Gln Pro Phe Gln Gln Phe Gly Arg Asp Val Ser Asp Phe Thr Asp
 545 550 555 560
 Ser Val Arg Asp Pro Lys Thr Ser Glu Ile Leu Asp Ile Ser Pro Cys
 565 570 575
 Ser Phe Gly Gly Val Ser Val Ile Thr Pro Gly Thr Asn Ala Ser Ser
 580 585 590
 Glu Val Ala Val Leu Tyr Gln Asp Val Asn Cys Thr Asp Val Ser Thr
 595 600 605
 Ala Ile His Ala Asp Gln Leu Thr Pro Ala Trp Arg Ile Tyr Ser Thr
 610 615 620
 Gly Asn Asn Val Phe Gln Thr Gln Ala Gly Cys Leu Ile Gly Ala Glu
 625 630 635 640
 His Val Asp Thr Ser Tyr Glu Cys Asp Ile Pro Ile Gly Ala Gly Ile
 645 650 655
 Cys Ala Ser Tyr His Thr Val Ser Leu Leu Arg Ser Thr Ser Gln Lys
 660 665 670
 Ser Ile Val Ala Tyr Thr Met Ser Leu Gly Ala Asp Ser Ser Ile Ala
 675 680 685
 Tyr Ser Asn Asn Thr Ile Ala Ile Pro Thr Asn Phe Ser Ile Ser Ile
 690 695 700
 Thr Thr Glu Val Met Pro Val Ser Met Ala Lys Thr Ser Val Asp Cys
 705 710 715 720
 Asn Met Tyr Ile Cys Gly Asp Ser Thr Glu Cys Ala Asn Leu Leu Leu
 725 730 735
 Gln Tyr Gly Ser Phe Cys Thr Gln Leu Asn Arg Ala Leu Ser Gly Ile
 740 745 750
 Ala Ala Glu Gln Asp Arg Asn Thr Arg Glu Val Phe Ala Gln Val Lys
 755 760 765
 Gln Met Tyr Lys Thr Pro Thr Leu Lys Tyr Phe Gly Gly Phe Asn Phe

770	775	780
Ser 785	Gln Ile Leu Pro Asp 790	Pro Leu Lys Pro Thr 795 Lys Arg Ser Phe Ile 800
Glu	Asp Leu Leu Phe 805	Asn Lys Val Thr 810 Leu Ala Asp Ala Gly Phe 815 Met
Lys	Gln Tyr Gly 820	Glu Cys Leu Gly Asp 825 Ile Asn Ala Arg Asp 830 Leu Ile
Cys	Ala Gln 835	Lys Phe Asn Gly 840 Leu Thr Val Leu Pro Pro 845 Leu Leu Thr
Asp 850	Asp Met Ile Ala Ala Tyr 855	Thr Ala Ala Leu Val Ser Gly Thr Ala
Thr 865	Ala Gly Trp Thr Phe 870	Gly Ala Gly Ala 875 Leu Gln Ile Pro Phe 880
Ala	Met Gln Met Ala 885	Tyr Arg Phe Asn Gly 890 Ile Gly Val Thr Gln 895 Asn
Val	Leu Tyr Gly 900	Asn Gln Lys Gln Ile 905 Ala Asn Gln Phe Asn 910 Lys Ala
Ile	Ser Gln 915	Ile Gln Glu Ser Leu 920 Thr Thr Thr Ser Thr 925 Ala Leu Gly
Lys	Leu 930	Gln Asp Val Val Asn 935 Gln Asn Ala Gln 940 Ala Leu Asn Thr Leu
Val 945	Lys Gln Leu Ser Ser 950	Asn Phe Gly Ala Ile 955 Ser Ser Val Leu Asn 960
Asp	Ile Leu Ser Arg 965	Leu Asp Lys Val Glu 970 Ala Glu Val Gln Ile 975 Asp
Arg	Leu Ile Thr 980	Gly Arg Leu Gln Ser 985 Leu Gln Thr Tyr Val 990 Thr Gln
Gln	Leu Ile 995	Arg Ala Ala Glu Ile 1000 Arg Ala Ser Ala Asn 1005 Leu Ala Ala
Thr	Lys 1010	Met Ser Glu Cys Val 1015 Leu Gly Gln Ser Lys 1020 Arg Val Asp
Phe	Cys 1025	Gly Lys Gly Tyr His 1030 Leu Met Ser Phe Pro 1035 Gln Ala Ala
Pro	His 1040	Gly Val Val Phe Leu 1045 His Val Thr Tyr Val 1050 Pro Ser Gln

Glu Arg Asn Phe Thr Thr Ala Pro Ala Ile Cys His Glu Gly Lys
 1055 1060 1065
 Ala Tyr Phe Pro Arg Glu Gly Val Phe Val Phe Asn Gly Thr Ser
 1070 1075 1080
 Trp Phe Ile Thr Gln Arg Asn Phe Phe Ser Pro Gln Ile Ile Thr
 1085 1090 1095
 Thr Asp Asn Thr Phe Val Ser Gly Asn Cys Asp Val Val Ile Gly
 1100 1105 1110
 Ile Ile Asn Asn Thr Val Tyr Asp Pro Leu Gln Pro Glu Leu Asp
 1115 1120 1125
 Ser Phe Lys Glu Glu Leu Asp Lys Tyr Phe Lys Asn His Thr Ser
 1130 1135 1140
 Pro Asp Val Asp Leu Gly Asp Ile Ser Gly Ile Asn Ala Ser Val
 1145 1150 1155
 Val Asn Ile Gln Lys Glu Ile Asp Arg Leu Asn Glu Val Ala Lys
 1160 1165 1170
 Asn Leu Asn Glu Ser Leu Ile Asp Leu Gln Glu Leu Gly Lys Tyr
 1175 1180 1185
 Glu Gln Tyr Ile Lys Trp Pro Trp Tyr Val Trp Leu Gly Phe Ile
 1190 1195 1200
 Ala Gly Leu Ile Ala Ile Val Met Val Thr Ile Leu Leu Cys Cys
 1205 1210 1215
 Met Thr Ser Cys Cys Ser Cys Leu Lys Gly Ala Cys Ser Cys Gly
 1220 1225 1230
 Ser Cys Cys Lys Phe Asp Glu Asp Asp Ser Glu Pro Val Leu Lys
 1235 1240 1245
 Gly Val Lys Leu His Tyr Thr
 1250 1255

<210> 50
 <211> 35
 <212> PRT
 <213> SARS coronavirus Urbani

<400> 50

Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln
 1 5 10 15

Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln
20 25 30

Leu Ser Ser
35

<210> 51
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 51

Ile Gln Ala Ala Leu Thr Lys Thr Ser Ala Ala Leu Gly Lys Leu Gln
1 5 10 15

Ala Ala Val Asn Arg Asn Ala Ala Ala Leu Asn Lys Leu Val Lys Ala
20 25 30

Leu Ser Ser
35

<210> 52
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>
<221> MISC_FEATURE
<222> (1)..(35)
<223> X=aminoisobutyric acid

<400> 52

Ile Gln Glu Ser Leu Thr Xaa Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15

Asp Val Val Asn Xaa Asn Ala Gln Ala Leu Asn Xaa Leu Val Lys Gln
20 25 30

Leu Ser Ser
35

<210> 53
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>
<221> MISC_FEATURE

<222> (1)..(35)
<223> X=dipropyl or dibutyl glycine

<400> 53

Ile Gln Glu Ser Leu Thr Xaa Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15

Asp Val Val Asn Xaa Asn Ala Gln Ala Leu Asn Xaa Leu Val Lys Gln
20 25 30

Leu Ser Ser
35

<210> 54
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>
<221> MISC_FEATURE
<222> (17)..(21)
<223> i,i+4 lactam bridge

<400> 54

Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15

Glu Val Val Asn Lys Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln
20 25 30

Leu Ser Ser
35

<210> 55
<211> 35
<212> PRT
<213> Artificial

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<220>
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<222> (7)..(11)
<223> i, i+4 lactam bridge

<220>
<221> MISC_FEATURE
<222> (28)..(32)
<223> i, i+4 lactam bridge

<400> 55

Ile Gln Glu Ser Leu Thr Glu Thr Ser Thr Lys Leu Gly Lys Leu Gln
1 5 10 15

Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Glu Leu Val Lys Lys
20 25 30

Leu Ser Ser
35

<210> 56
<211> 35
<212> PRT
<213> Artificial

<220>
<223> synthetic peptide

<220>
<221> MISC_FEATURE
<222> (14)..(21)
<223> i, i+7 bridge

<400> 56

Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Glu Leu Gln
1 5 10 15

Asp Val Val Asn Glu Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln
20 25 30

Leu Ser Ser
35

<210> 57
<211> 35
<212> PRT
<213> Artificial

<220>
<223> synthetic peptide

<400> 57

Ile Ile Glu Ser Leu Thr Thr Thr Ile Thr Ala Leu Gly Lys Leu Ile
1 5 10 15

Asp Val Leu Asn Gln Asn Ile Gln Ala Leu Asn Thr Leu Ile Lys Gln
20 25 30

Leu Ser Ser
35

<210> 58
<211> 35
<212> PRT
<213> SARS coronavirus Urbani

<400> 58

Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile Asp
1 5 10 15

Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp Leu
20 25 30

Gln Glu Leu
35

<210> 59
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 59

Ile Ala Ala Ile Asn Lys Ser Val Ala Ala Ile Gln Lys Glu Ile Ala
1 5 10 15

Arg Leu Asn Glu Val Ala Lys Ala Leu Asn Ala Ser Leu Ile Arg Leu
20 25 30

Gln Ala Leu
35

<210> 60
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>
<221> MISC_FEATURE
<222> (1)..(35)
<223> X=aminoisobutyric acid (Aib)

<400> 60

Ile Ser Gly Ile Asn Xaa Ser Val Val Asn Ile Gln Lys Glu Ile Asp
1 5 10 15

Arg Leu Asn Xaa Val Ala Lys Asn Leu Asn Xaa Ser Leu Ile Asp Leu
20 25 30

Gln Glu Leu
35

<210> 61
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>

<221> MISC_FEATURE
<222> (1)..(35)
<223> X=dipropyl or dibutyl glycine

<400> 61

Ile Ser Gly Ile Asn Xaa Ser Val Val Asn Ile Gln Lys Glu Ile Asp
1 5 10 15

Arg Leu Asn Xaa Val Ala Lys Asn Leu Asn Xaa Ser Leu Ile Asp Leu
20 25 30

Gln Glu Leu
35

<210> 62
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>
<221> MISC_FEATURE
<222> (16)..(20)
<223> i,i+4 lactam bridge

<400> 62

Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile Glu
1 5 10 15

Arg Leu Asn Lys Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp Leu
20 25 30

Gln Glu Leu
35

<210> 63
<211> 35
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>
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<222> (6)..(10)
<223> i,i+4 lactam bridge

<220>
<221> MISC_FEATURE
<222> (27)..(31)
<223> i,i+4 lactam bridge

<400> 63

Ile Ser Gly Ile Asn Glu Ser Val Val Lys Ile Gln Lys Glu Ile Asp
1 5 10 15

Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Lys Leu
20 25 30

Gln Glu Leu
35

<210> 64
<211> 35
<212> PRT
<213> Artificial

<220>
<223> synthetic peptide

<220>
<221> MISC_FEATURE
<222> (13)..(20)
<223> i,i+7 bridge

<400> 64

Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Glu Glu Ile Asp
1 5 10 15

Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp Leu
20 25 30

Gln Glu Leu
35

<210> 65
<211> 35
<212> PRT
<213> Artificial

<220>
<223> synthetic peptide

<400> 65

Ile Ser Gly Ile Asn Ala Ser Ile Val Asn Ile Gln Lys Glu Ile Asp
1 5 10 15

Arg Leu Asn Glu Val Ile Lys Asn Leu Asn Glu Ser Leu Ile Asp Leu
20 25 30

Gln Glu Leu
35

<210> 66
<211> 39
<212> PRT
<213> SARS coronavirus Urbani

<400> 66

Asp Leu Gly Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln
1 5 10 15

Lys Glu Ile Asp Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser
20 25 30

Leu Ile Asp Leu Gln Glu Leu
35

<210> 67
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 67

Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile
1 5 10 15

Asp Arg Leu Asn Glu Val Ile Lys Asn Leu Asn Glu Ser Leu Ile Asp
20 25 30

Leu Gln Glu Leu
35

<210> 68
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 68

Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile
1 5 10 15

Ala Arg Leu Asn Glu Val Ala Lys Ala Leu Asn Glu Ser Leu Ile Asp
20 25 30

Leu Gln Glu Leu
35

<210> 69
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 69

Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile
1 5 10 15

Ala Arg Leu Asn Glu Val Ile Lys Ala Leu Asn Glu Ser Leu Ile Asp
Page 37

20

25

30

Leu Gln Glu Leu
35

<210> 70
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 70

Asp Ile Ala Ala Ile Asn Ala Ser Val Ala Asn Ile Gln Lys Glu Ile
1 5 10 15

Ala Arg Leu Asn Glu Val Ala Lys Ala Leu Asn Glu Ser Leu Ala Ala
20 25 30

Leu Gln Ala Leu
35

<210> 71
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>
<221> MISC_FEATURE
<222> (17)..(21)
<223> relative to residues 1166 to 1170; lactam bridge

<400> 71

Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile
1 5 10 15

Glu Arg Leu Asn Lys Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp
20 25 30

Leu Gln Glu Leu
35

<210> 72
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<220>
<221> MISC_FEATURE
<222> (17)..(21)

<223> salt bridge

<400> 72

Asp Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile
1 5 10 15

Glu Arg Leu Asn Lys Val Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp
20 25 30

Leu Gln Glu Leu
35

<210> 73

<211> 36

<212> PRT

<213> Artificial

<220>

<223> Synthetic peptide

<400> 73

Asp Ile Glu Glu Ile Asn Lys Lys Val Glu Glu Ile Gln Lys Lys Ile
1 5 10 15

Glu Glu Leu Asn Lys Lys Ala Glu Glu Leu Asn Lys Lys Leu Glu Glu
20 25 30

Leu Gln Lys Lys
35

<210> 74

<211> 36

<212> PRT

<213> Artificial

<220>

<223> Synthetic peptide

<220>

<221> MISC_FEATURE

<222> (1)..(36)

<223> Introduction of salt bridges relating to mutations departing from
SEQ ID NO:48 (HR-C4a extended; 1150-1185)

<400> 74

Asp Ile Ser Gly Ile Asn Ala Ser Val Val Glu Ile Gln Lys Lys Ile
1 5 10 15

Glu Glu Leu Asn Lys Lys Ala Glu Glu Leu Asn Lys Lys Leu Ile Asp
20 25 30

Leu Gln Glu Leu
35

<210> 75

<211> 7

<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 75
Ile Gln Glu Ser Leu Thr Thr
1 5

<210> 76
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 76
Thr Ser Thr Ala Leu Gly Lys
1 5

<210> 77
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 77
Leu Gln Asp Val Val Asn Gln
1 5

<210> 78
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 78
Asn Ala Gln Ala Leu Asn Thr
1 5

<210> 79
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 79
Leu Val Lys Gln Leu Ser Ser
1 5

<210> 80

<211> 14
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 80

Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys
1 5 10

<210> 81
<211> 14
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 81

Thr Ser Thr Ala Leu Gly Lys Leu Gln Asp Val Val Asn Gln
1 5 10

<210> 82
<211> 14
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 82

Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr
1 5 10

<210> 83
<211> 14
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 83

Asn Ala Gln Ala Leu Asn Thr Leu Val Lys Gln Leu Ser Ser
1 5 10

<210> 84
<211> 21
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 84

Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15

Asp Val Val Asn Gln
20

<210> 85
<211> 21
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 85

Thr Ser Thr Ala Leu Gly Lys Leu Gln Asp Val Val Asn Gln Asn Ala
1 5 10 15

Gln Ala Leu Asn Thr
20

<210> 86
<211> 21
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 86

Leu Gln Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr Leu Val
1 5 10 15

Lys Gln Leu Ser Ser
20

<210> 87
<211> 28
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 87

Ile Gln Glu Ser Leu Thr Thr Thr Ser Thr Ala Leu Gly Lys Leu Gln
1 5 10 15

Asp Val Val Asn Gln Asn Ala Gln Ala Leu Asn Thr
20 25

<210> 88
<211> 28
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 88

Thr Ser Thr Ala Leu Gly Lys Leu Gln Asp Val Val Asn Gln Asn Ala
Page 42

1 5 10 15

Gln Ala Leu Asn Thr Leu Val Lys Gln Leu Ser Ser
20 25

<210> 89
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 89

Ile Ser Gly Ile Asn Ala Ser
1 5

<210> 90
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 90

Val Val Asn Ile Gln Lys Glu
1 5

<210> 91
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 91

Ile Asp Arg Leu Asn Glu Val
1 5

<210> 92
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic peptide

<400> 92

Ala Lys Asn Leu Asn Glu Ser
1 5

<210> 93
<211> 7
<212> PRT
<213> Artificial

<220>

<223> Synthetic peptide

<400> 93

Leu Ile Asp Leu Gln Glu Leu
1 5

<210> 94

<211> 14

<212> PRT

<213> Artificial

<220>

<223> Synthetic peptide

<400> 94

Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu
1 5 10

<210> 95

<211> 14

<212> PRT

<213> Artificial

<220>

<223> Synthetic peptide

<400> 95

Val Val Asn Ile Gln Lys Glu Ile Asp Arg Leu Asn Glu Val
1 5 10

<210> 96

<211> 14

<212> PRT

<213> Artificial

<220>

<223> Synthetic peptide

<400> 96

Ile Asp Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser
1 5 10

<210> 97

<211> 14

<212> PRT

<213> Artificial

<220>

<223> Synthetic peptide

<400> 97

Ala Lys Asn Leu Asn Glu Ser Leu Ile Asp Leu Gln Glu Leu
1 5 10

<210> 98

<211> 21

<212> PRT

<213> Artificial

<220>

<223> synthetic peptide

<400> 98

Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile Asp
1 5 10 15

Arg Leu Asn Glu Val
20

<210> 99

<211> 21

<212> PRT

<213> Artificial

<220>

<223> synthetic peptide

<400> 99

Val Val Asn Ile Gln Lys Glu Ile Asp Arg Leu Asn Glu Val Ala Lys
1 5 10 15

Asn Leu Asn Glu Ser
20

<210> 100

<211> 21

<212> PRT

<213> Artificial

<220>

<223> synthetic peptide

<400> 100

Ile Asp Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser Leu Ile
1 5 10 15

Asp Leu Gln Glu Leu
20

<210> 101

<211> 28

<212> PRT

<213> Artificial

<220>

<223> synthetic peptide

<400> 101

Ile Ser Gly Ile Asn Ala Ser Val Val Asn Ile Gln Lys Glu Ile Asp
1 5 10 15

Arg Leu Asn Glu Val Ala Lys Asn Leu Asn Glu Ser
20 25

<210> 102

<211> 28

<212> PRT
<213> Artificial

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Applicant: Regents of the University of Colorado
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1. Certificate of Mailing by Express Mail - 1 page
2. PCT International Application Transmittal Letter - 1 page
3. PCT Request in paper form - 4 pages
4. Fee Calculation page in paper form - 1 page
5. Specification excluding sequence listing (description - 63 pages; claims - 5 pages; abstract - 1 page) - 69 pages
6. 23 sheets of drawings
7. Sequence listing; paper - 46 pages
8. Sequence listing; Computer Readable Form/CRF - 1 diskette
9. Statement under 37 CFR 1.821-824 - 1 page
10. PCT-Safe file on diskette (3-05WOX.zip; 3-05WOX.log)
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